



SEQUENCE LISTING

<110> Miller, Brian S.
Shetty, Jayarama K.

<120> Modified Forms of Pullulanase

<130> GC396-2

<140> 09/262,126

<141> 1999-03-03

<160> 9

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 2794

<212> DNA

<213> B. deramificans

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 <223> Xaa = Any Amino Acid

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 35 40 45
 Pro Trp Ser Leu Trp Met Trp Pro Lys Asp Gly Gly Gly Ala Glu Tyr
 50 55 60
 Asp Phe Asn Gln Pro Ala Asp Ser Phe Gly Ala Val Ala Ser Ala Asp
 65 70 75 80
 Ile Pro Gly Asn Pro Ser Gln Val Gly Ile Ile Val Arg Thr Gln Asp
 85 90 95
 Trp Thr Lys Asp Val Ser Ala Asp Arg Tyr Ile Asp Leu Ser Lys Gly
 100 105 110
 Asn Glu Val Trp Leu Val Glu Gly Asn Ser Gln Ile Phe Tyr Asn Glu
 115 120 125
 Lys Asp Ala Glu Asp Ala Ala Lys Pro Ala Val Ser Asn Ala Tyr Leu
 130 135 140
 Asp Ala Ser Asn Gln Val Leu Val Lys Leu Ser Gln Pro Leu Thr Leu
 145 150 155 160
 Gly Glu Gly Xaa Ser Gly Phe Thr Val His Asp Asp Thr Ala Asn Lys
 165 170 175
 Asp Ile Pro Val Thr Ser Val Lys Asp Ala Ser Leu Gly Gln Asp Val
 180 185 190
 Thr Ala Val Leu Ala Gly Thr Phe Gln His Ile Phe Gly Gly Ser Asp
 195 200 205
 Trp Ala Pro Asp Asn His Ser Thr Leu Leu Lys Lys Val Thr Asn Asn

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Leu Tyr Gln Phe Ser Gly Asp Leu Pro Glu Gly Asn Tyr Gln Tyr Lys				
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Val Ala Leu Asn Asp Ser Trp Asn Asn Ser Tyr Pro Ser Asp Asn Ile				
		245		250
Asn Leu Thr Val Pro Ala Gly Gly Ala His Val Thr Phe Ser Tyr Ile				
		260		265
Pro Ser Thr His Ala Val Tyr Asp Thr Ile Asn Asn Pro Asn Ala Asp				
		275		280
Leu Gln Val Glu Ser Gly Val Lys Thr Asp Leu Val Thr Val Thr Leu				
		290		295
Gly Glu Asp Pro Asp Val Ser His Thr Leu Ser Ile Gln Thr Asp Gly				
305		310		315
Tyr Gln Ala Lys Gln Val Ile Pro Arg Asn Val Leu Asn Ser Ser Gln				
		325		330
Tyr Tyr Tyr Ser Gly Asp Asp Leu Gly Asn Thr Tyr Thr Gln Lys Ala				
		340		345
Thr Thr Phe Lys Val Trp Ala Pro Thr Ser Thr Gln Val Asn Val Leu				
		355		360
Leu Tyr Asp Ser Ala Thr Gly Ser Val Thr Lys Ile Val Pro Met Thr				
		370		375
Ala Ser Gly His Gly Val Trp Glu Ala Thr Val Asn Gln Asn Leu Glu				
385		390		395
Asn Trp Tyr Tyr Met Tyr Glu Val Thr Gly Gln Gly Ser Thr Arg Thr				
		405		410
Ala Val Asp Pro Tyr Ala Thr Ala Ile Ala Pro Asn Gly Thr Arg Gly				
		420		425
Met Ile Val Asp Leu Ala Lys Thr Asp Pro Ala Gly Trp Asn Ser Asp				
		435		440
Lys His Ile Thr Pro Lys Asn Ile Glu Asp Glu Val Ile Tyr Glu Met				
		450		455
Asp Val Arg Asp Phe Ser Ile Asp Pro Asn Ser Gly Met Lys Asn Lys				
465		470		475
Gly Lys Tyr Leu Ala Leu Thr Glu Lys Gly Thr Lys Gly Pro Asp Asn				
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Val Lys Thr Gly Ile Asp Ser Leu Lys Gln Leu Gly Ile Thr His Val				
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Gln Leu Met Pro Val Phe Ala Ser Asn Ser Val Asp Glu Thr Asp Pro				
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Thr Gln Asp Asn Trp Gly Tyr Asp Pro Arg Asn Tyr Asp Val Pro Glu				
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Gly Gln Tyr Ala Thr Asn Ala Asn Gly Asn Ala Arg Ile Lys Glu Phe				
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Lys Glu Met Val Leu Ser Leu His Arg Glu His Ile Gly Val Asn Met				
		565		570
Asp Val Val Tyr Asn His Thr Phe Ala Thr Gln Ile Ser Asp Phe Asp				
		580		585
Lys Ile Val Pro Glu Tyr Tyr Tyr Arg Thr Met Ile Gln Val Ile Ile				
		595		600
Pro Thr Asp Gln Val Leu Glu Met Lys Leu Xaa Ala Glu Arg Pro Met				
		610		615
Val Gln Lys Phe Ile Ile Asp Ser Leu Lys Tyr Trp Val Asn Glu Tyr				
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His Ile Asp Gly Phe Arg Phe Asp Leu Met Ala Leu Leu Gly Lys Asp				
		645		650
Thr Met Ser Lys Ala Ala Ser Glu Leu His Ala Ile Asn Pro Gly Ile				
		660		665
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Ala	Leu	Tyr	Gly	Glu	Pro	Trp	Thr	Gly	Gly	Thr	Ser	Ala	Leu	Pro	Asp	
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	690					695					700					
Phe	Asn	Asp	Asn	Leu	Arg	Asn	Ala	Leu	Asp	Gly	Asn	Val	Phe	Asp	Ser	
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Ser	Ala	Gln	Gly	Phe	Ala	Thr	Gly	Ala	Thr	Gly	Leu	Thr	Asp	Ala	Ile	
			725						730					735		
Lys	Asn	Gly	Val	Glu	Gly	Ser	Ile	Asn	Asp	Phe	Thr	Ser	Ser	Pro	Gly	
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Glu	Thr	Ile	Asn	Tyr	Val	Thr	Ser	His	Asp	Asn	Tyr	Thr	Leu	Trp	Asp	
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Lys	Ile	Ala	Leu	Ser	Asn	Pro	Asn	Asp	Ser	Glu	Ala	Asp	Arg	Ile	Lys	
	770					775					780					
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Lys	Ala	Gln	Tyr	Pro	Asp	Val	Phe	Asn	Tyr	Tyr	Ser	Gly	Leu	Ile	His	
	835						840					845				
Leu	Arg	Leu	Asp	His	Pro	Ala	Phe	Arg	Met	Thr	Thr	Ala	Asn	Glu	Ile	
	850					855					860					
Asn	Ser	His	Leu	Gln	Phe	Leu	Asn	Ser	Pro	Glu	Asn	Thr	Val	Ala	Tyr	
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Glu	Leu	Thr	Asp	His	Val	Asn	Lys	Asp	Lys	Trp	Gly	Asn	Ile	Ile	Val	
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Val	Tyr	Asn	Pro	Asn	Lys	Thr	Val	Ala	Thr	Ile	Asn	Leu	Pro	Ser	Gly	
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Lys	Trp	Ala	Ile	Asn	Ala	Thr	Ser	Gly	Lys	Val	Gly	Glu	Ser	Thr	Leu	
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His	Pro	Val	Thr	Phe	Gly	Lys	Ile	His	Cys	Val	Arg	Ala	Ser	Ser	Gly	
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Thr	Val	Asp	Gln	Tyr	Ala	Lys	Ala	Val	Thr	Val	Asn	Gly	Glu	Lys	Gly		
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Val	Val	Leu	Arg	Pro	Asp	Gln	Met	Lys	Trp	Thr	Ala	Pro	Leu	Lys	Pro		
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Phe	Ser	His	Pro	Val	Asp	Ala	Val	Ile	Tyr	Glu	Thr	His	Leu	Arg	Asp		
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Ala	Leu	Thr	Glu	Thr	Asp	Thr	Gln	Thr	Ala	Asn	Gly	Ser	Ser	Ser	Gly		
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Phe	Trp	Asp	Lys	Met	Ser	Phe	Ala	Leu	Pro	Gln	Glu	Asn	Asp	Ser	Arg		
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Trp	Asp	Arg	Arg	Glu	Thr	Phe	Lys	Glu	Asp	Val	His	Tyr	Ile	Arg	Arg		
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	610					615					620						
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Ile	Ala	Tyr	Arg	Leu	Tyr	Asp	Leu	Asp	Glu	Val	Asp	Glu	Trp	Lys	Asp		
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Glu	Asp	Pro	Thr	Glu	Ile	Lys	Lys	Thr	Val	Ala	Val	Asn	Gly	Ile	Gly		
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Pro	Asp	Thr	Pro	Asp	Asn	Gln	Asp	Val	Val	Val	Arg	Leu	Pro	Asp	Val		
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Ala	Val	Pro	Gly	Glu	Ala	Val	Thr	Ala	Val	Glu	Asn	Gln	Ala	Val	Ile		
	50					55					60						
His	Leu	Val	Asp	Ile	Ala	Gly	Ile	Thr	Ser	Ser	Ser	Ala	Ala	Asp	Tyr		
65				70					75					80			
Ser	Ser	Lys	Asn	Leu	Tyr	Leu	Trp	Asn	Asn	Glu	Thr	Cys	Asp	Ala	Leu		
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Ser	Ala	Pro	Val	Ala	Asp	Trp	Asn	Asp	Val	Ser	Thr	Thr	Pro	Ser	Gly		
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Ser	Asp	Lys	Tyr	Gly	Pro	Tyr	Trp	Val	Ile	Pro	Leu	Asn	Lys	Glu	Ser		
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Gly	Cys	Ile	Asn	Val	Ile	Val	Arg	Asp	Gly	Thr	Asp	Lys	Leu	Ile	Asp		
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Ser	Asp	Leu	Arg	Val	Ala	Phe	Gly	Asp	Phe	Thr	Asp	Arg	Thr	Val	Ser		
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Arg	Ala	Ala	Phe	Gly	Val	Ala	Leu	Ala	Glu	Ala	His	Trp	Val	Asp	Lys		
			180					185					190				
Asn	Thr	Leu	Leu	Trp	Pro	Gly	Gly	Gln	Asp	Lys	Pro	Ile	Val	Arg	Leu		
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Tyr	Tyr	Ser	His	Ser	Ser	Lys	Val	Ala	Ala	Asp	Gly	Glu	Gly	Lys	Phe		
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Thr	Asp	Arg	Tyr	Leu	Lys	Leu	Thr	Pro	Thr	Thr	Val	Ser	Gln	Gln	Val		
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Ser	Met	Arg	Phe	Pro	His	Leu	Ser	Ser	Tyr	Ala	Ala	Phe	Lys	Leu	Pro		

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Ser	Asp	Lys	Val	Ala	Asp	Ile	Gln	Gln	Pro	Phe	Ser	Arg	Leu	Cys	Glu																																	
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Trp	Thr	Thr	Asp	Tyr</																																												

Leu	Asn	Pro	Asp	Ile	Tyr	Phe	Phe	Gly	Glu	Gly	Trp	Asp	Ser	Asn	Gln	705	710	715	720
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Glu	Ile	Asp	Tyr	Asn	Gly	Ala	Pro	Gly	Gly	Tyr	Ala	Ala	Asp	Pro	Thr	820	825	830	
Glu	Val	Val	Asn	Tyr	Val	Ser	Lys	His	Asp	Asn	Gln	Thr	Leu	Trp	Asp	835	840	845	
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Pro	Glu	Ser	Arg	Thr	Leu	Asn	Glu	Phe	Ala	Gly	Glu	Thr	Leu	Gln	Leu	1025	1030	1035	104
Ser	Ala	Ile	Gln	Gln	Thr	Ala	Gly	Glu	Asn	Ser	Leu	Ala	Asn	Gly	Val	1045	1050	1055	
Gln	Ile	Ala	Ala	Asp	Gly	Thr	Val	Thr	Leu	Pro	Ala	Trp	Ser	Val	Ala	1060	1065	1070	
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